



**Robust ensembles  
for Assessment and  
scenario study**

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S. Galmarini

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# A science-based use of ensembles of opportunities for assessment and scenario study: a re-analysis of HTAP-1 ensemble

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## Abstract

The multi-model ensemble exercise performed within the HTAP project context (Fiore et al., 2009) is used here as an example of how a *pre-inspection*, diagnosis and selection of an ensemble, can produce much better and more reliable results. This procedure is contrasted with the often-used practice of simply averaging model simulations, assuming model difference as equivalent to independence, and using the diversity of simulation as an illusory estimate of model uncertainty. It is further and more importantly demonstrated how conclusions can drastically change when future emission scenarios are analysed using an un-inspected ensemble. The HTAP multi-model ensemble analysis is only taken as an example of a wide spread and common practice in air quality modelling.

## 1 Introduction

A multi-model (MM) ensemble is defined as a group of simulations of the same case study, produced by formally different models, which are statistically treated in an attempt to improve the quality of the result (Potempski and Galmarini, 2009). Given the ever increasing collaborations of geophysical modelling communities in joint assessment studies, MM ensembles are becoming very popular and an opportunity to extend and generalize individual deterministic model results (Solazzo et al., 2012, 2014; Solazzo and Galmarini, 2014; Galmarini et al., 2004; Vautard et al., 2012; Evans et al., 2013; Bishop and Abramowitz, 2013).

In particular in atmospheric sciences, MM ensembles are used extensively in climate and air quality predictions and assessments. While in climate research and applications many of the concepts applied and described here are well known and correctly used, in air quality this is not always the case and several are the examples of direct use of *un-inspected* ensembles. We shall describe an *inspected* ensemble (opposed to an un-inspected one) as: a set of model results, whose properties and characteristics,

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which could lead to coinciding (diverging) biased results and a false sense of agreement (uncertainty).

MM ensembles derived from simply different models are said to be potentially prone to redundancy and overconfidence. The inspection is therefore primarily finalised at:

- the identification of the level of diversity (communality) shared by the model results,
- retaining only those that are contributing with original information
- removing the redundancy.

Techniques exist that allow such screenings that rely on the existence of observations and the comparison of the ensemble variability with the observational variability (Potemski and Galmarini, 2009; Solazzo et al., 2013).

In this study we aim at demonstrating the importance of using existing good practices in the air quality MM ensemble context. Toward the scope we have selected a case study published in the past which does not exploit the true value of having multiple model results at hand. The case analyzed is the HTAP (Hemispheric Transport of Air Pollution) phase 1 multi-model exercise (Dentener et al., 2010) and in particular the multi-model ensemble activity performed within it and presented by Fiore et al. (2009). The study of Fiore et al. (2009) is used here as mere representative of a wide spread practices in the air quality modelling communities at all scales and it represents just an example on how things could be improved further. An additional reason for selecting this case is the fact that ensembles are used by Fiore et al. (2009) for sensitivity studies with respect to emission reduction options. The inspection of the ensemble can have important consequences also for emission scenarios as shown later, an aspect never considered before in the literature.

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is always a better result than that of one model. That would be true if the models were independent but there is no a-priori proof of that. Some questions arise: how robust are the results if the models are not independent models? How different would be the result should some model not take part to the activity or more outliers like the one present in the Fig. 1 would be present? How generalised is the result since the selection of the ensemble members is based on the voluntary participation to a joint activity and the ensemble does not contain all possible results? Is there any duplication of information? Is all the information contained in a MM ensemble relevant and necessary? Since the construction of ensemble is not governed by scientific selection criteria, so it happens that the subsequent ensemble result strictly depends on *aleatory* factors and one can presume that it lacks generality.

The screening methodology proposed and that we will apply as an example to the FetA09 set, is a good way to exploit an abundance of model results in the best way, to transform the aleatory gathering of information into a more robust result that is based on scientific principles. The large ensemble of model results becomes an opportunity to *cherry-pick* those models that produce the most accurate MM ensemble and use only those to drive conclusions. The analysis will help identifying the size of the non-redundant ensemble and the subsets of members to produce skilled results.

### 2.1 Inspecting a multi model ensemble

In this section the MM ensemble of FetA09 is inspected. We will concentrate on the ozone simulations over the same regions presented in FetA09 and we will make use of exactly the same model data and observations used by Fiore et al. (2009). The inspection is based on the following steps:

- determine to what extent the variability present in the observation is reproduced by the ensemble,
- determine the minimum number of models necessary to represent the observed variability,

- identification of the models that will be part of the reduced ensemble which will be subsequently used.

### 2.1.1 The “accounted” variability: eigen-analysis and ranked histogram technique

5 The goal of this first analysis is to determine to what extent the observational variability is reproduced by the ensemble. An optimal situation is the one in which the variability of observations coincides with that produced by the ensemble of models, in other words the ensemble of the results all together covers the same range of variation of the measurements. Any deviation from this condition, namely a smaller or a larger variability of the ensemble with respect to the observed one would show, on one side, the incapacity of the ensemble to model the observed reality, or on the other, the addition of irrelevant information to the simulation of the observed situation. Therefore considering that a MM ensemble is assembled on an opportunity basis rather than results characteristics, this first step is of primary importance to estimate to what extent the gathered set is appropriate for the case study.

15 A technique to assess the variability and to estimate the redundancy of the MM ensemble with respect to that of the observations, was suggested by Annan and Hargreaves (2010) and applied in some MM ensemble modelling contexts (see, e.g. Solazzo et al., 2013; Solazzo and Galmarini, 2014). It consists of projecting the observation anomalies (the element-wise difference between the observations and their mean) onto the principal components (PCs) of the covariance matrix of the deviation of the ensemble of models from the MM mean (the element-wise difference between each model realisation and the MM ensemble mean). Principal component analysis (Jolliffe, 2002) is probably the most well-known and wide-spread dimension-reduction technique. It is based on eigen-analysis to select uncorrelated directions associated with the largest variances.

25 When applied to the HTAP 21-member ensemble analysed by FetA09, this method shows that the first (largest) eigenvalue already explains more than 90 % of the ob-

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servational variability in most regions, the only exception being Japan with 60 %. In other words, most of the ensemble members have a significant projection onto the first eigen-vector defining the major component, thus explaining the same portion of variance. If too many models are projected on the same eigenvector, it means that there are too many models producing repeating or “overlapping” solutions (thus, the ensemble is redundant). A well-behaved MM ensemble (not necessarily the theoretical case of independent models) should be made of a number of models whose eigenvalues contribute to the explanation of as many different components as the observational variability and the ratio model-to-observed variance should be close to unity. In the case of the HTAP MM ensemble, when all eigen-values are taken into account, the MM ensemble variance is 4.7, 6.0, 8.7 times the variance of the observation anomalies for the EU Mediterranean, Central 0–1 km, and Central 1–2 km, regions respectively. Concerning the US Mountains, Great Lakes, SE, NE, SW regions, the full MM ensemble mean accounts for 25.4, 9.1, 20.6, 10.7, 5.6 times the observed variability, respectively, and finally 4.7 times for the Japanese sub-region. According to the definition of Annan and Hargreaves (2010) the ensemble is therefore *wide*, i.e. its variability is larger than the observed one. Dealing with a wide ensemble implies that there is a substantial amount of redundant variability, i.e. variability already accounted for by other models. Not all information contained in the ensemble is needed in principle and needs to be reduced.

An alternative method to diagnose the variability spanned by an ensemble of models to the eigenvalues used is the Talagrand or Ranked Histogram (RH) (Talagrand et al., 1998), which provides an evaluation of the consistency of the ensemble with an observed quantity. In a RH the observations are ranked into a number of bins equal to the number of models making up the ensemble plus one for the extremes. The ensemble members are sorted to define ranges or “bins” of the modeled variable such that the probability of occurrence of the observation within each bin is, ideally, equal. The bins are determined by ranking the ensemble members from lowest to highest. The interval between each pair of ranked values forms a bin. If there are  $N$  ensemble members,

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then there will be  $N + 1$  bins (Hamill, 2001). The underlying assumption is that each ensemble member in principle can introduce an independent degree of variability. An indication of an ill-constructed ensemble is the ratio between the number of elements and the number of data available per model. If there are  $N$  models with time series each of size  $n_t$  (elements of the time series), the implication of  $N > n_t$  is that there will be at least  $N - n_t$  empty bins in the RH, indicating redundancy of the ensemble and that the ensemble is inappropriate for the case analyzed. This same result could be visualized by looking at the load factors resulting from the decomposition in PCs: many projections would be null, as the number of Eigen-vector is larger than the number of data to project. The HTAP MM ensemble used in this example,  $N = 21$  and  $n_t = 12$ . The RH for the nine sub-regions is reported in Fig. 2. Six (NA NE) to nine (NA SW) bins out of 21 are populated, (i.e. contain non-zero values), due to insufficient data and excess of redundant information. The use of the Ranked histogram reveals another important problem with the FetA09 ensemble. Good ensemble practice would require  $n_t \gg N$ . The plots clearly show that there are many empty bins (so degrees of freedom in the process that are not part of the reality as no observations are present in that range). The uneven distribution of the histograms shows that much emphasis (overconfidence) is given to some aspects of the process description, while others are neglected, that is another way of representing the redundancy obtained with PC analysis presented earlier.

### 2.1.2 Effective number of models

Having assessed that the ensemble is redundant it is important to determine the minimum number of models from those available in the ensemble that would suffice to describe the observational variability. A very robust method never used in air quality is that developed by Bretherton et al. (1999). The effective number of models sufficient to

reproduce the variability of the observation is defined as:

$$N_{\text{eff}} = \frac{\left(\sum_{k=1}^N \lambda_k\right)^2}{\sum_{k=1}^N \lambda_k^2} \quad (1)$$

with  $\lambda$  eigenvalue of the **corr**( $d_i, d_j$ ) matrix, which contains the linear correlation coefficient between any pair  $d_i, d_j$  ( $i, j = 1, \dots, N$ ).  $d$  is a metric defined accordingly to Pennel and Reichler (2011):

$$d_m = e_m - RMME \quad (2)$$

where the index  $m$  identifies the model, MME is the multi model error (the average of all individual model's errors) and  $R$  is the Pearson correlation coefficient between  $e_m$ , the error of model  $m$  and the MME. The removal of MME in Eq. (2) makes model errors more dissimilar from one another and uncovers "hidden" trends that are outweighed by overarching commonalities. Indeed the scope of the metric  $d_m$  is to determine similarities between models beyond the dominating ones induced by shared inputs and/or common parameterisations to the extent that the former are accounted for in the average. Expression (1) should be interpreted as: only if all eigenvalues were equal to unity, Eq. (1) would take a value of  $N_{\text{eff}} = N$ , which corresponds to the situation where all directions are equally important and all models add independent contributions to the explanation of the observational variability. On the other hand, if all error fields were similar, only one eigenvalue would be non-zero and  $N_{\text{eff}} = 1$ . Equation (1) provides an analytical estimate of the dimensions of the subspace of models necessary to produce the information of the whole ensemble.

For the HTAP MM ensemble of FetA09, Eq. (1) gives  $N_{\text{eff}}$  ranging between  $\sim 2$  and 4 for the regions analysed by FetA09 compared to the original 21 models. Thus, approximately three quarter of the available information on variance is redundant. This is a very revealing result that indicates paradigmatically the relevance of a pre-inspection

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of an ensemble. What seemed like a largely populated ensemble turns out to be incapable of capturing several degrees of freedom of observations and 2 to 4 members of 21 are sufficient to describe the observational variability. One may ask: if so, why is the average of the 21 models fitting so well with the observations as presented in FetA09?

The answers could be: pure chance, since finally the model results participated out of good will, and happened to be there in the right mixture. Just consider what would have happened to the mean of the models should one of the two most evident outliers in Fig. 1 decide to withdraw from the exercise. Alternatively an explanation could be the massive smoothing due to the monthly averaging along with the high level of tuning of the models around specific solutions that are normally distributed around the average observed data.

### 2.1.3 Reducing ensembles

As demonstrated in the previous sections, the HTAP MM ensemble is redundant and in particular 2 to 4 members are sufficient to represent the observational variability while the rest do not add any new information. Similarly, the extra elements are likely to deteriorate any evaluation metrics applied to the ensemble. At this point we know that the number of models that are necessary and sufficient is smaller than 21 but we do not know which combination of members for every grouping produces the optimal ensemble.

Given  $N$  members, there are  $G = N!/[r!(N-r)!]$  possible groups of  $r$  elements. A straight forward way to identify the optimal ensemble (optimal sub set) and maximize the accuracy of the ensemble is to analyse all the  $G$  combinations of subsets of models and identify the one that minimize the Root Mean Square Error (RMSE). The latter is a measure of the accuracy (the even distribution of model results around the observed value), and high accuracy also improves precision (a reduced spread/scatter of the model results around the observed value). In fact while accuracy is a pre-requisite for precision, the contrary does not hold.

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In Fig. 3 we report the curves of minimum, mean, and maximum RMSE for the nine sub-regions used by FetA09 as a function of the number of members of ensembles ( $r = 2, \dots, 21$ ). The figure confirms the results on the number of models necessary to maximize the ensemble performance and tells us that which combination of the 2 to 4 models out of 21 produces such improvement. The scores of the reduced ensemble are reported in Table 2 and are compared against the ones produced by the full ensemble mean. In all cases the mean of the reduced ensemble improves the accuracy (from 31 % for NA NW to 71 % for NA Mountain and NA Lakes) and precision (most notably for NA SE and NA NE). As it can be seen in several regions the use of the full ensemble of opportunity produces a clear deterioration in the ensemble statistics. In Table 2 we report also the ranking of the models contributing to minimize the error in the sub-regions. As from the table it is often the case that the error is minimized by mix-ranked (good performing and bad performing) of members. In fact, if the two best models have a high chance of being also highly correlated then they would share some portion of information thus resulting redundant. Therefore when considering the ensemble mean of these two models, very little decrease in error would be found compared to the individual models. Mathematically, the theorems by Elashoff et al. (1967) and Cover (1974) have proven two important results on the selection of member and evaluation of individual scores: the best two models are seldom the combination of two models that maximises the score of an ensemble average, and furthermore, that the best single model may not appear in the ensemble maximising the feature score. As a result, the simple method of making ranked combinations of models with the best individual features may prove unsuccessful, as also demonstrated by e.g. Solazzo et al. (2013), Hannan and Hargreaves (2011), and others. This confirms the importance of the inspection of the available results prior to their use and of having at disposal a large pool of models from which optimal subsets can be extracted.











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**Table 1.** Number of effective models  $N_{\text{eff}}$  for the sub-regions object of the analysis (with reference to Fig. 2 of Fiore et al., 2009, top panel, based on  $\text{corr}(d_i, d_j)$ ). nrec is the number of surface receptors used for evaluation.

Sub-region	$N_{\text{eff}}$
EU Mediterranean region (nrec = 6)	4.0
EU central region 0–1 km (nrec = 24)	3.1
EU central region 1–2 km (nrec = 11)	3.5
NE-USA (nrec = 13)	1.9
SW USA (nrec = 5)	1.8
SE USA (nrec = 6)	1.9
Great Lakes USA (nrec = 8)	2.0
Mountainous USA (nrec = 10)	1.8
Japan EANET (nrec = 10)	2.6

**Table 2.** RMSE-ranking and scores of the reduced MM ensemble mean for the sub-regions object of the analysis (RMSE: Root-Mean-Square-Error; PCC: Pearson Correlation Coefficient;  $\sigma$ : ratio of the modelled to the observed SD).

Domain	Ranking of the MinRMSE combination	score
EU central 0–1 km	1, 15, 19	RMSE = 1.69 (2.65) PCC = 0.98 (0.96) $\sigma$ = 0.99 (1.10)
EU central 1–2 km	7, 17, 18	RMSE = 3.35 (9.2) PCC = 0.98 (0.95) $\sigma$ = 1.03 (1.25)
EU medit	4, 6, 13, 15, 19	RMSE = 0.76 (1.44) PCC = 0.99 (0.98) $\sigma$ = 1.0 (1.13)
NA SW	8, 10, 11, 15	RMSE = 2.0 (2.9) PCC = 0.95 (0.96) $\sigma$ = 0.87 (0.86)
NA SE	1, 2, 4, 8	RMSE = 3.61 (10.27) PCC = 0.77 (0.62) $\sigma$ = 0.83 (1.81)
NA NE	3, 5, 6, 7	RMSE = 3.01 (7.8) PCC = 0.93 (0.90) $\sigma$ = 0.90 (1.56)
NA Mountain	1, 5, 12	RMSE = 1.53 (5.33) PCC = 0.93 (0.90) $\sigma$ = 1.04 (1.44)
NA Lakes	1, 5, 6	RMSE = 1.89 (6.58) PCC = 0.97 (0.91) $\sigma$ = 1.03 (1.45)
Japan EANET	12, 15	RMSE = 3.11 (5.70) PCC = 0.96 (0.79) $\sigma$ = 0.66 (0.51)

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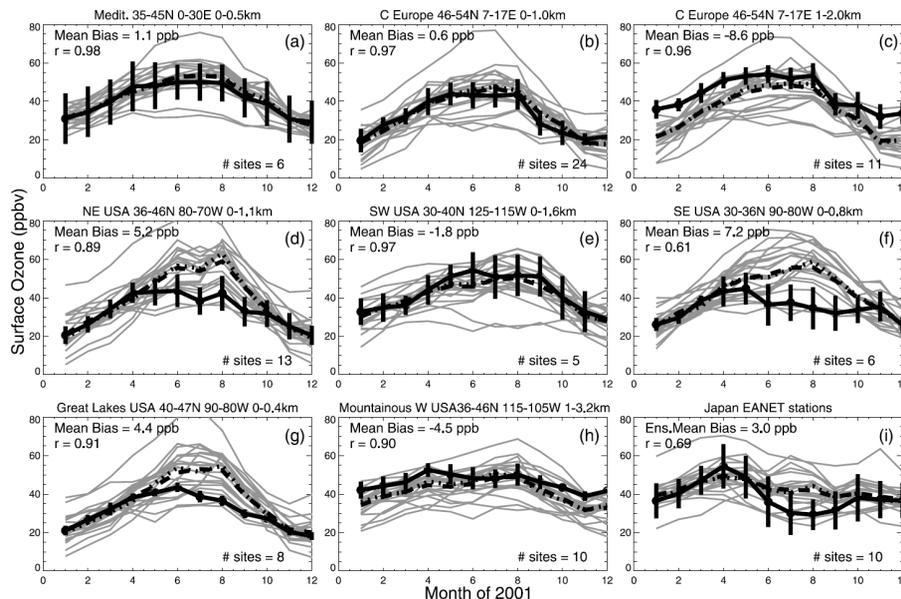
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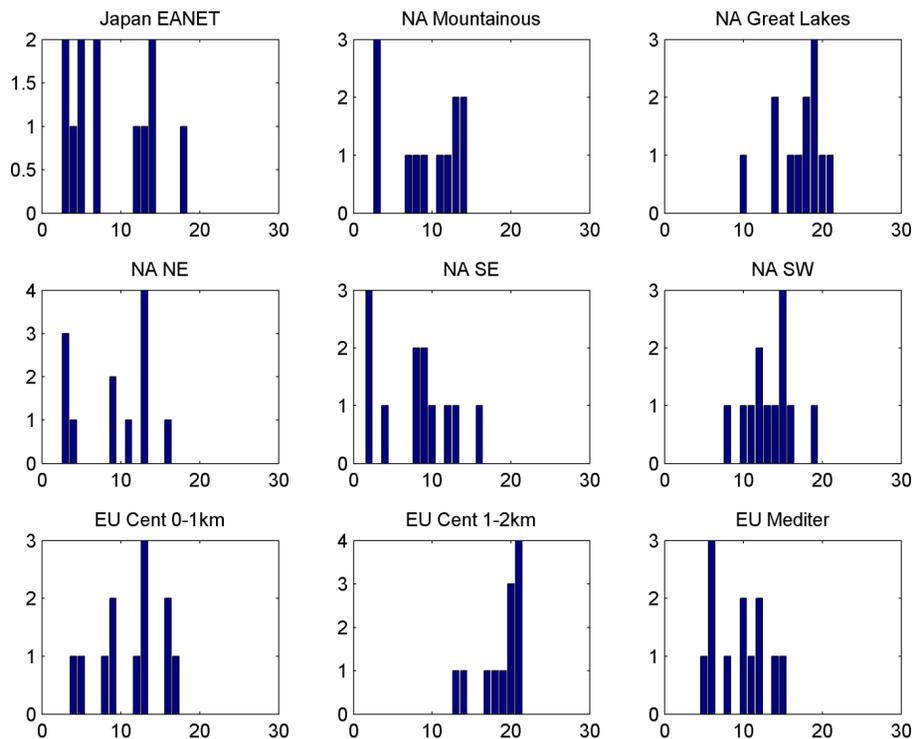
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**Figure 1.** From Fiore et al. (2009): monthly mean surface  $O_3$  concentrations (ppb) for the year 2001. Observed values (black circles) represent the average of all sites falling within the given latitude, longitude, and altitude boundaries and denoted by the symbols in Fig. 1; vertical black lines depict the SD across the sites. Monthly mean  $O_3$  in the surface layer of the SR1 simulations from the 21 models are first sampled at the model grid cells containing the observational sites and then averaged within subregions (gray lines); these spatial averages from each model are used to determine the multimodel ensemble median (black dotted line) and mean (black dashed line). Observations are from CASTNET (<http://www.epa.gov/castnet/>) in the United States, from EMEP (<http://www.nilu.no/projects/ccc/emepdata.html>) in Europe, and from EANET (<http://www.eanet.asia/>) in Japan.

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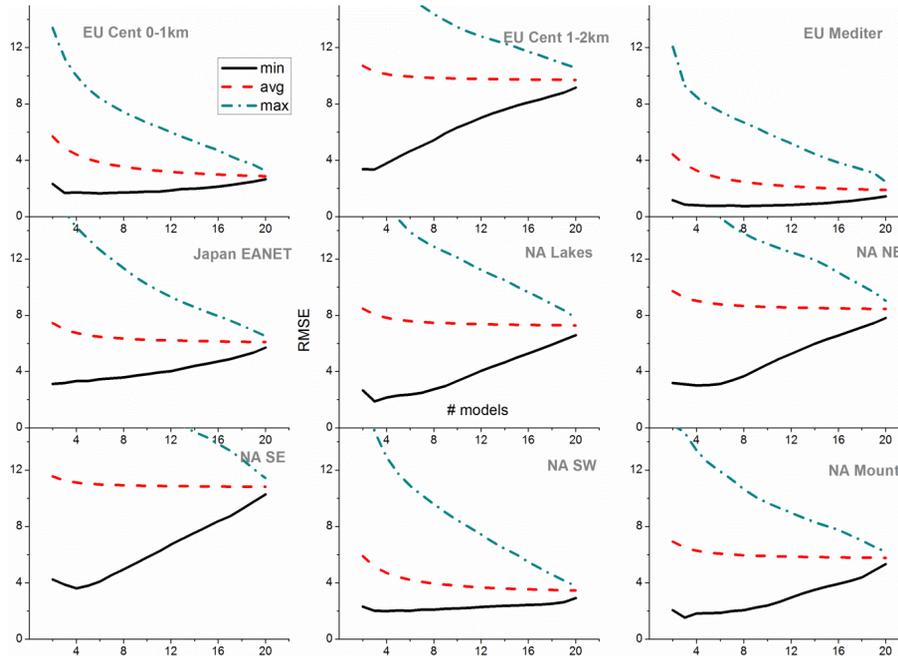
**Figure 2.** Ranked histogram for the nine sub-regions subject to MM ensemble evaluation.

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**Figure 3.** Maximum (dash-dot), average (dashed), and minimum (continuous line) RMSE for all subsets of MM combinations and for the nine sub-regions subject to MM ensemble evaluation.

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